130

Glu Glu Met Lys Lys His Glu Ser Asn Asn Val Gly Lou Leu Glu Asn 428 Leu Thr Asn Gly Val Thr Ala Gly Asn Gly Asp Asn Gly Leu Ile Pro 440 Gln Arg Lys Ser Arg Thr Pro Glu Asn Gln Gln Phe Pro Asp Asn Glu 455 460 Ser Glu Glu Tyr His Arg Ile Cys Glu Leu Val Ser Asp Tyr Lys Glu 465 470 478 Lys Gln Met Pro Lys Tyr Ser Ser Glu Asn Ser Asn Pro Glu Gln Asp 485 490 495 Leu Lys Leu Thr Ser Glu Glu Glu Ser Gln Arg Leu Glu Gly Ser Glu 500 Asn Gly Gin Pro Glu Leu Glu Asn Phe Met Als Ile Glu Glu Met Lys S15 S20 S25 Lys His Gly Ser Thr His Val Gly Phe Pro Glu Asn Leu Thr Asn Gly 335 Ala Thr Ala Gly Asn Gly Asp Asp Gly Leu Ile Pro Pro Arg Lys Ser 545 550 555 Arg Thr Pro Glu Ser Gin Gln Phe Pro Asp Thr Glu Asn Glu Glu Tyr 568 570 575 His Ser Asp Glu Gln Asn Asp Thr Gln Lys Gln Phe Cys Glu Glu Gln 588 Asn Thr Gly Ile Lew His Asp Glu Ile Lew Ile His Glu Glu Lys Gln 595 600 The Glu Val Val Glu Lys Met Asn Ser Glu Leu Ser Leu Ser Cys Lys S1S 615 620 Lys Glu Lys Asp Ile Leu His Glu Asn Ser Thr Leu Arg Glu Glu Ile 629 630 £38 640 Ala Met Leu Ary Leu Glu Leu Asp Thr Met Lys His Gln Ser Gln Leu 645 850

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<2115 671

<212> PRT

<213> Homo sapien

<400> 380

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Leu	asp	Arg 195	Arg	Cys	gin	Leu	Asn 200	Val.	Leu	Asp	Asn	Lys 205	Lys	Arg	The
Ala	Leu 210	ïle	Lys	Ala	Val	Gln 215	Сув	Gln	Glu	Asp	320 220	Cys	Ala	Leu	Met
225					Thr 230					235					240
				245	Ala				250					255	
			260		Gly			268					270		
Leu	Thr	Pro 275	Leu	2,60	Leu	Gly	7&1 280	His	Glu	Gln	Lys	Gln 285	Gln	Val	Val
	290				Lys	299					300				
305					11e				•	325	•				320
				325	Glu				330					335	
			340		Arg			345					350		
		385			Ser		380					365			
	370				Assi	375			-		389				
385					290 390					395					400
				405	Glu				410					415	
			420		His			425					0.8.9		
		435			Thr		440					445			
	450			•	Thr	455					460				
865	ಚಾತ್ರಾಣ	ಟಾಖ.ಟ.	7.À.T.	F2 1 28	Arg 470	33.5	೧೩೫	w.c.u	ಬಜಚ	475	262	ນສຄ	a ya	A. J. S.	480
	Gln	Met	Pro	Lys 485	Tyr	Sex	Ser	Glu	Asn 490		Asn	Pro	@lu	01n 495	
₽₩	Lys	Leu	Thr 500	Ser	Glu	Glu		Ser SOS	Gln	Arg	ren	Gju	Gly 510		Sla
Asn	Gly	Gla S1S	Fro	Glu	Lys	Arg	Ser S20	Gln	\$1u)ro	Slu	Ile 525	Asn	Lys	Asp
Gly	Asp 530	Arg	Glu	Leu	Glu	Asn 535	Fhe	Met	Ala	Ile	Glu 540	Glu	Mest	Lys	Fåå
	Gly	Ser	Thr	Mis	Val	Gly	Phe	Pro	Glu		Leu	Thr	Assi	Gly	
545	8 % 2	A 4	×	M 5	550	N		2	V 5	555	85 and 15	W	*	89.000	860
				565	Asp				570					578	
			580		Gin Asp			595					596		
ris ris Tr	oxes go	595	19:44.6	31.05.14	wali	.2.025	600	m k to	MALES.	e-moste	ir I, is	605	للة بدائن	13.44.6	.ex.25.15
Thr	Gly 610		Leu	His	Asp	Glu 615	ile	Leu	lle	His	61u 620	Ğlu	Lys	Gin	Ile
Glu	val	val.	Glu	Lys	zeM	neA	Ser	Glu	Leu	Sex	Leu	Ser	Cys	Lys	Lys

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635
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                                         538
                                                              640
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cotragitoto tecentroas iceatosico atologosico agigggicat telgatoasi 660
quantiques tarreagese tgescarges cetesurges tecceuntge cetegaguagg 720
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geatectyca gatygicocy geocicates tyctyacety tetycayyya etytectect 840
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gaccigigct biotygigig gagicosggg cigciaggsa aaggaatggg cagacacagg 1440
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acacacages aggrigacge igisaacsis geocacgeig teeiggggge acigggaage 1740
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ggggaagggt cccttttgcm tigccmagtg ccmlasccmt gagcmctmct ctaccmtggt 180
totgootoot ggoomagomag yotggittyo mmagamtymma tymmigatto tacayotmagy 240
acttaacctt gasatggaaa gictigcaat occatiligea ggatoogist gigcacatgs 300
chotytagag agoagoatto coagggacot tygaaaacagt tygcactyta agytycttyc 360
tococcaagac acatoctasa aggigitgia aiggigaasa ogicitocit cittatigec 420
cottottatt talgigaaca acigitigic bilittigia tottibitaa acigiaaagt 480
traattytya aastyssatat ratyraasta asttatyrys tittititte aasytasaas 540
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totoasagoo atotgotgto ttogagtaog gacacateat vactootgoa ttgttgatca 180
amacytygay ytyottttoo tomyotmaym myonottayo mamayotoym mtagaottay 240
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gogacottgg ocogsagget otagosaggs ecoacogace coagoogog oggoggedge 180
yoşşaciliş ocoşşiştiyi gşşşoşşago ggacişoştş toogoşgacş şşcaşoşaaş 240
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coccetering tyreatcaty atragracet atgagetrzyg casasagetre threagagge 120
tgaaccagga coggettetg ggoggetgaa aggggcaagg aggcaaggac cocgtetete 180
ccacggatgg ggagagggca ggaggagacc cagccaagtg cottttootc agcactgagg 240
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goggerrage actionicag acacaactic theotypics iccaginging gygatcaica 360
cttacccscc coccssqttc asgaccssat cttccsgctg coccettogt gtttccctgt 420
gtttgrigta grigggratg irtoraggaa oraagaagor otragrotigg tgtagtrico 480
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gttigaagat igootoitot acagoitotg agaatigigt taltitosott goosagigaa 180
ggaccocctc occaacatge occagedcar coctaageat ggtocettgt caccaggeaa 240
ccaggaaact gotactigtg gacctcacca gagaccagga gagtttggtt agctcacagg 300
acticoccoa occoagaaga tiagcatocc atactagact catactoaac tcaactaggc 360
tratartras tigalggita tisquraatt cratitotti riggitalis taaaragasa 420
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aacgactttc caaataatct caccagogoc ttocagotca ggogtoctag aagogtottg 180
associates cossiste tigistico totoaccos distocicae ascisasant 240
cocaggazac etteagacta cottectety cettoageza yyggegttye eracattete 300
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getetangay tetganenga ntegttysee cantntgaca naaggaaagg eggagettat 180
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tagocagggo actgotgoca acagocagto conataccat catginacco ggigogotot 180
naantingat niccanaged diadecaten tagitelget didecadegg niaccageed 240
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ttgccgggaa cactgcagag acaatgctgt gagtttccaa cottagccca totgcgggca 180
qaqaaqqtot aqtitqtoca toaqoattat catqatatoa qqaotqqtta otiqqttaaq 240
gaggggtota ggagabotgt coottitaga qacacottac ttataatgaa gtattiggga 300
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cathtattas testecetge eigtgtetat tattatatte atatetetae getggasaet 420
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                                                                   566
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graggaggas egggetttaa ggagttttaa gotgagtgte aetgtagase esaaatassa 188
torcaagatt atogggagaa agggggdagt aattaccoaa atooggttyy agoatgacyt 240
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ccasctactt stotscaatt statcticaa gaataccots sccatocott tsactsacst 300
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teactactyt geotegacca gtgaggagag ctggacegae agegaggtgg acteateatg 180
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coggeatiga gogdatggge cogstygges togaccacat ggsstocans attgansgca 240
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tataccetet caccatecec tigictaete tgaigecece aagaigeaae igggeageia 360
gttgycccca taattetygg cettigttyi tigttitaat taetigggea teecaggaag 420
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<211> 355
<212> DNA
<213> Homo sapiens
<220%
<221> misc_feature
<222> (1)...(355)
<223> n = A.T.C or G
<400> 401
acceptance temperate acceptants are acceptage and the second acceptance acceptant and acceptance acceptant acceptance acc
tgatgtoton aagtagtona oottoattta actotttgaa actgtatoat ottigooaag 120
taagagiggt ggockattto agotgottty acasaatgae tygotootga oktaacgike 180
tataaatgaa tytyotigaay caaaytyooc atyytyycyy cyaagaagan asagatytyt 240
tttgttttgg actototgtg gtocottoca atgotgmggg tttocaacca ggggaagggt 360
```

```
contrition traceaugty coataseest gagesetant classatogn totac
                                                                   355
<210> 402
<2115 407
<212> DNA
<213> Nomo sapiens
<2200
<221> misc_feature
<222> (1)...(407)
<223> n * A,T,C or G
<400> 402
atggggcaag etggataaag aaccaagace cactggagta tgetgtette aagaaaccca 60
totoacatgo ggtggcatac ataggotoaa aataaaxggaa tggagaaaaa tatttoaago 120
aaatggaaaa cagaaaaaag caggtgttgc actcctactt totgacaaaa cagactatgc 180
gastamagat maaamagaga aggacattac amaggtggtc otgaccittg atmestetca 240
ttgcttgata ccascctggg ctgttttaat tgcccaascc aasaggstas tttgctgagg 300
ttgtggaget teteccetge agagagtece tgateteera aaatttggtt gagatgtaag 360
gnigatititg cigacaacic cilitotgaa gittiacica titocaa
<210> 403
<2110 303
<212> DNA
<213> Homo sapiens
e2200
<221> misc_feature
<222> (1)...(303)
<223> n * A,T,C or G
<400> 403
caytatitat ayonmaarty maaaqotmyt ayongyraay totraaator ayyoncomam 60-
tootaagoaa gagooatggo atggtgaasa tgcaaaagga gagtotggoo aatotacaaa 120
tagagaacaa gacctactca gtcatgaaca aaaaggcaga caccaacatg gatctcatgg 180
gggattggat attgtaatta tagagcagga agatgacagt gatcgtcatt tggcacaaca 240
tettaacaae gaeegaaace cattatttae ataaacetee atteggtaae catgttgaaa 300
gga
<210> 404
<211> 225
<212× DNA
<213> Homo sapiens
<400> 404
aagtytaact tttassaatt tagtygatti tyssaaatict tagaggaaag tasaggaasa 60
attgitaatg cactcatita cetitacatg gigaaagite tetetigate etacaaacag 120
acattiticoa chogigitto catagitgit aagigiatoa gaigigitgg goatgigaat 180
ctoraagigo oigigiaata aataaagiai oittattica ticat
                                                                   225
<230> 405
<233> 334
<212> DWA
<213> Homo sapiens
<220>
<221> misc_feature
<222> (1)...(334)
```

```
<223 × n × A,T,C or G
<400> 405
gagotyttat actylgagit chactagyaa atcalcaaat cigagygity toigyaggac 60
thoastacac checoccest agigasteag officeagggg giccagiocc telecitaet 120
teateresat cocatgoraa aggaagaroo terstootig gotraragoo tistotaggo 180
ttoccagtgo etocaggaca gagtgggtta tgttttcago tocatocttg otgtgagtgt 240
ctggtgegg: tgtgeeteea gettetgete agtgetteat ggacagtgte cageecatgt 300
cartotocae betetranny tygatrocae cort
                                                                   3.34
<230> 496
<2315 216
<212> DNA
<213> Homo sapiens
<2220×
<221> miso_fmature
<222> (1)...(216)
<223> n * A,T,C or G
<400> 406
tttcatacct aatgagggag ttganatnac atnnsaccag gaaatgcatg gatctcaang 60
gasacasada cocaatasad toggastago asastaadaa cigtgagada tgcactigot 120
sensaaeses aattinaigi tgeaecetig titetaeaee igigggitat gacaaagaes 180
actgocasag astriticasg asggaggact gocant
62305 407
<211> 413
<212> DNA
<213> Homo sapiens
<400> 407
gotgacttgc tagtatcatc tgcattcatt gaagcacaag aacttcatgc ctigactcat 60
gtasatgcas taggattasa asatasattt gatatoacat ggasacagac asasasatatt 120
gtacaacatt gcacocagig teagaiteta cacciggera etcaggaage aagagttaat 180
cocagaggto tatgloctas tgtgttatgg casalggatg toatgcacgt accticatit 240
ggassatigi catitytoca tytyscagti gatactiati cacalitical atygycaacc 300
tgccagacag gagaaagtot teecatgita asagacatti attatetigi titeetgica 360
tqqqaqttcc aqaaasaqtt aaaacagaca atqqqccagg ttctqtagta aag
<210> 408
<211x 183
<212> DNA
<213> Romo sapiens
<330×
<221> misc_feature
<222> (1)...(183)
<2235 n = A.T.C or G
<400> 408
ggagetngee eteaatteet eeathtetat gttancatat ttaatgtett tignnattaa 60
tnottaacta gttaatoott aaagggotan ntaatootta actagtooot coattgtgag 120
cattateett eeagtateen eestetnitt tattiactee tieetggeta eecatgiact 180
nee.
                                                                   183
<210> 409
<211> 250
```

```
<212> DMA
<213> Homo sapiens
<220×
<221> misc feature
<222> (1)...(250)
\langle 223 \rangle n = A,T,C or G
<400> 409
cocacguatg staagetett tatttetgta agtretgeta ggaaateate aaatetgaeg 60
gtggtbtggg ggacctgaac aaacctootg taattaatca gcbbtcagtt totoccccta 120
gtocctcott caacaacata ggaggatoot cocottcttt etgeteaegg cettatetag 180
gotteccagt goodcaagga dagegtigggd tatgtitada gegontoott gottgiggigig 240
ggcontatge
<210> 410
×211× 306
<23.25 DNA
<213> Romo sapiens
<2200
<221> misc_feature
<222> (1)...(306)
<223 \times n \approx A,T,C \text{ or }G
<400> 410
ggotggtttg caaqaatgaa atgaatgatt otacagotag qaottaacot tgaaatggaa 60
agtictigosa toccattigo aggateogto igigoscatg coictigiaga gagoagcatt 120
cocagggace tiggaaacag tiggeacigt aaggigetig eiceceaaga cacateetaa 180
asggigtigt aalggigaas seegelleet telttalige eestlettat tlatgigase 240
nactygttyg ettititityn atettittita aactygaaag ticaattyng aaaatyaata 300
tentge
<210> 411
<211> 261
<212> DWA
<213> Homo sapiens
<228×
<221> misc_feature
<222> (1)...(261)
<223> n = A,T,C or G
<400> 411
agagatatin citagginaa agittoataga gittoccatga actatatgac iggccacaca 60
ggalottitg tattiaagga itoigagati tigciigago aggatiagat aaggoigito 120
titasaigic tgaasiggaa cagatiicas aassasacce cacaatctag ggigggaaca 180
aggaaggaaa gatgigaata ggoigaiggg caaaaaacca attiacccai cagiiccagc 260
ottototosa ggngaggcas a
c2105 412
<211> 241
<212> DNR
<213 > Homo sapiens
<220×
<221> misc_feature
<222> (1)...(241)
```

```
\langle 223 \rangle n = A.T.C or G
<400> 412
gttmastgtt acctgacatt totacaacac occactoace gatgtattog tigoccagig 80
ygaacatace agootgaatt tggaaaaaat aattgtyttt ottyccoagg aaatactacy 120
actgaettig atggeteese sascatasee esgigissas sesgasgaig iggaggggag 180
cigggagatt tracigggta cattgaatto ccasactaco cangcastta occasocaso 240
<210> 413
<2115 231
<212> DNA
<213> Homo sapiens
<220>
<221> misc_feature
<222> (1)...(231)
<223> n * A,T,C or G
<400> 413
machettaca atomasgtgm ctcatctgtg tgcttgmatc ctttccactg tctcatctcc 60
ctcatccaag titctagtac ctictctitg tigtgaagga taalcaaact gaacaacaa 120
aagtitacto teeteatitig gaacetaaaa actetettet teetigigtet gagggeteea 180
agastrotty astrantict cayatratty gygacarcan atraggasor t
<23.0> 43.4
<211> 234
<212> DWA
<213> Homo sapiens
<400> 414
actificate asgraction cagasoring approaches carragates tracegoese for
gatggagotg aaaacataac coactotyte otyggaggcac tygggaagoot agagaaggot 120
gtgagocsag gagggagggt cttrotttgg catgggatgg ggatgaagta aggagaggga 189 🕟
etggacecce tggaagetga tteactatgg ggggaggtgt attgaagtee teea
<210> 415
<211> 217
<212> DNA
<213> Homo sapiens
€220>
<221> misc_feature
<222> (1)...(217)
<223 > n * A.T.C or G
<400> 415
grataggatt aagactgagt atottitota cattottita actitotaag gggcacttot 60
caassoscag accaggiage assicteds tgototaagg nictoscosc cartitotos 120
cacctagosa tagtagaatt cagtootact totgaggoos gaagaatggt toagassaat 180
antggattat aaasaatsac aattaagaaa aataatc
<210> 416
<211> 213
<212> DWA
<213> Homo sapiens
<220>
```

```
<221> misc_festure
<222× (1)...(213)
<223> n = A,T,C or G
<400> 436
atgearatht assignment geotogothe tagasgacat diggnotion otoligostica 60
ggcacagcag tasagctott tgatteccag astcaagaac toteccette agactattse 120
cgaatgcaag gtggttaatt gaaggccact aattgatget caaatagaag gatattgact 180
atattggaac agatggagto totactacaa aag
<210> 417
<211> 303
<232> DNA
<213> Nomo sapiens
<220>
<221> misc_feature
<222> (1)...(303)
<223 > n = A,T,C or G
<400> 417
nagiciticag goccaticagg gaagiticaca ciggagagaa gicatacata igiacigist 60
qtqqqaaaqq ctitactctq agttcaaatc ttcaagccca tcagagagtc cacactggag 120
agaagocata caaatgcaat gagtgtggga agagcitcag gagggattoo cattatcaag 180
tteatetagt ggteeacses ggsgagaast cetatssatg tgsgatatgt gggaaggget 240
trantrasag thogtstott casatrosto ngasggnora cagtatanan assocttita 300
agt
<210> 418
<211> 328
<212> DWA
<213> Homo sapiens
₹220%
<221> misc_feature
<222> (1)...(328)
<223> n = A,T,C or G
<400> 418
tititggogg tggtggggca gggacgggac angagtotca cicigitgcc caggotggag 60
tgcacaggea tgatetegge teactacaac coetgectee catgteeaag egattettgt 120
gootoagoot toootgtago tagaattada ggdadatgoo accadacooa gdtagttttt 180
gtattitiag tagagacagy gitteaceat ghiggeragg eiggleicaa acteeinaee 240
tragnggtra ggrtggtote asactoriga ortraagiga teigoprace trageriere 390
assgtgctan gattacaggc egtgagcc
                                                                   328
<210> 419
<231> 389
<212> DNA
<213> Homo sapiens
<2220×
<221> misc_feature
<222> (1)...(389)
<223> n = A,T,C or G
<400% 42.9
cotoctomag acgycotyty ytocycotoc cyycamocmm ymagoctycm ytycomiaty 60
```

```
accordage categactom agertgasag gragegtaca cortector gatertgety 120
cttgtttcct ctctgtggct coaltcatag cacagttgtt gcactgaggc ttgtgcaggc 180
cyagosaggo caagotyyst caaayagosa coagtosact otycoacygt ytyocayyos 240
coggetator agarancese attactaget congessatt gasastasgt tetterace 300
taaaggtagg accaaagggc atctgetttt cigaagteet etgetetate agecateaeg 360
tggcagccac tenggctgtg tcgacgcgg
<210> 420
<211> 408
c212> DWA
<213> Homo sapiens
<480> 420
gttoctecta actorigora gasacagoto toctoascat gagagotgoa corotoctec 60
tggccagggc agcaagectt agecttggct tettgtttet getttttte tggctagace 120
gaagtgtact agocaaggag tigaagtiig igaciiiggi giiinggcat ggagaccgaa 180
gtoccatigs cassities actgasses tasaggasts steatggers casggatitg 240
gocaactcan ocagotgggo atggagcago attatgaact tggagagtat ataagaaaga 300
gatatagaaa attotigaat gagtootata aacatgaaca ggtttatatt ogaagcacag 360
acyttgaccy gactitgaty aagtgetaty acasacctgy caageecy
<210> 421
<211> 382
<212> DNA
<213> Homo sapiens
<220×
<221> misc feature
<222> (1)...(352)
<223> n = A,T,C or G
<400> 421
gotoaaasat otitiitactg atmygostyy otacacaato attyactatt acygagyoca 60
gaggagaatg aggootggoo tgggagooot gtgootacta naagcacatt agattatoca 120
treactgacs gasesggtet ittitigggte circticies scesenatat actigosgie 190
ctectiotig asgaticiti ggesgrigic titgicatas cocsosgig tagasacasg 240
ggtgcascat gasatttetg tttegtages agtgestgte tesesagttg gesngtetge 300
cactorgagt trattgggtg titgtricct rigagators igratified gg
<210> 422
<211> 337
<212> 0WA
<213> Homo sapiens
<400> 422
stycraccat gotygosaty cagogygogy togsagyoot gratatocay occasgotyy 60
ogatyatoga oggosacogt bycocqaagt tycogatyon agocqaagog gtagtosagg 120
gogatagoaa ggigorggog alogoggogg ogtoaaloot ggocaaggio agcogigato 180
gtgaaatggc agctgtegaa ttgatetaec egggttatgg categgeggg cataaggget 240
atoogacaco ggtgcacotg gaagoottgo agoggotggg googacgeog attoacogac 300
gettetteeg eeggtaegge tggeetatga aaattat
                                                                  337
<210> 423
<211> 310
<212> DNA
<213> Homo sapiens
<330×
```

```
<221> misc_feature
e222> {1}...{310}
<223> n = A,T,C or G
<400> 423
getraasaat ettittaety atalygesty getseasaat caltgactat tagaggerag 60
aggagaatga ggootggoot gggagoootg tgootactam aagoncatta gattatocat 120
trangarag aaraggtott tilligggtot fictionea coargatata citigoagier 180
techticings againeting gesquister tignestase cesesgigt ansascasgg 340
gtggaacatg asatttotgt tiogtagoaa gigcatgtot cacagttgto aagtotgooc 300
teegagttta
                                                                   310
c210> 424
<211× 370
<212> DMA
<213> Homo sapiens
<2220>
<221> misc feature
<222> (1)...(370)
<223> n ~ A.T.C or G
<400> 424
gotosassat offittacty staggosigg clacacasto attgactatt agaggotaga 60
agassatgas gootgeoots saascotest sootactasa ascacattas attatocatt 120
cantgacaga acaggietht bitgggtoob teltobecad cacgatabae tigcagbeet 180
cettettgaa gattetttgg cagttgtett tgteataace cacaggtgta gasacatest 240
ggitgaatoi coiggaacto coicattagg talgasatag calgaigcal igsalaaagt 300
cacqaaqqtq qcasaqatca caacqctqcc caqqanaaca ttcattqtga taaqcaqqac 360
teegtegacg
<230> 425
<23.1> 23.6
<212> DNA
<213> Homo sapiens
e220>
<221> misc_feature
<222> (1)...(216)
<223 > n * A.T.C or G
<400> 425
sattgetatn ntttatttig cesetessa testiseess assassass intisastgs 60
taacaacnea acateaaggn aaananaaca ggaatggnig acintgcata aainggeega 120
anattateca itainitaag ggitgactic aggntacage acacagacaa acatgeceag 190
qaqqninica qqaccqcicq aiginiinig aqqaqq
<210> 426
<211> 596
<212> DNA
<213> Homo sapiens
<400> 426
officeagiga ggataaccet gtigococgg geogaggite tecattagge feigatigat 60
tygcagtoag tyatggaagg gtyttotgal cattocyact gooccaaggy togotygoca 120
gotototgtt tigotgagtt ggoagtagga octaattigt taattaagag tagatggtga 180
getgiortig tattitgatt aarotsatgy cottoorage argadioyga ticagotyga 240
garatrangg caarititaa igaaatgati igaagggrea tiaagaggea ritoregita 300
```

```
ttaggcagti catolgoact galaacttot tggcagotga gotggtogga gotgtggooc 360
asacgcaçaç tiggetittig gittigagat Adaactetia atetitiagi caigetigag 420
ggtggatggz ottbbbagct ttaacockat bigcactgcc biggaagtgt agccaggaga 480
atacantoat atactogigg gettagagge cacagoagat gicalitygic tactgootga $40
gtocogotgg toccatocca ggacettoca toggogagta cetgggagec cgtgct
<210> 427
<211> 107
<212> DNA
<213> Homo sapiens
<2205
<221> misc_feature
<222> (1)...(107)
<223 » n » A,T,C or G
<400> 427
gaagaattoa agtiaggitt attoasaggg ottschgaga atootahaco cagghoocag 60
congression goottamaga gotoctottt gactgooogg otdagng
<210> 428
<211> 39
c212> DMA
<213> Homo sapiens
<220×
<221> misc_feature
<222> (1)...(38)
<223> n = A,T,C or G
2400% 828
gaactteena ansangaett tattesetat titaeatt
                                                                   38
<2105 429
<211> 544
<212> DNA
<213> Homo sapiens
<400> 429
ctttgctgga oggaatasaa gtggaogcaa gcatgacctc otgatgaggg ogctgcattt 60
attgsagage ggetgeagee etgeggttea gattasaate egagaattgt atagaegeeg 120
atalocarga actoligasg gactitoliga titalocara alcasatoat eggillitoag 180
thiggaiggt ggotoatcac olytagaacc igactiggco giggoliggaa tocactogit 240
gestiesast teagitasas steasteass atestetest gttggttetg tgstgettea 300
agatactaag occacattig agaigcagoa gocatotoco coaattocto otgiocatoo 360
tgatgtgrag ttaasaaasto tgecotttta tgatgtoott gatgttotca toasgcoose 420
gagtttagit caaagcagta ttcagegatt tcaagagaag tttittattt ttgctttgac 480
acctmaacaa gitagagaga tatgmatatm magggattit tigmmaggig giaggagaga 548
3522
                                                                   544
<210> 430
<211> 507
<212> DNA
<213> Nomo sapiens
<220×
<221> miso feature
<222× (1)...(507)
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<223> n * A,T,C or G <400> 430 cttatonosa tyggyotoco saacttyyot ytycaytyya sactocyyyy yaattigas 60 gascactgac accestette cacceegaca etetgattta attgggetge agtgagaaca 120 gagoatcaat thamaaagot goodagaatg tintootggg dagogtigtg abottigeen 180 cottogigae titaigeaat geateatget alticatase taatgaggga gitocaggag 240 atteaserag gatgitteta encetytygg tiatgacasa gacaseigee aasgaatnii 300 caaqaaqqaq qactqcaaqt atatoqtqqt qqaqaaqaaq gacccaaaaa aqacctqttc 360 tytosytysa tyystastor astyryotto raytagycso aggyntocca yycosyyon 429 cattotocto tegecototaa tagtoaatga tigigiagoo algootatoa giaaaaaagat 480 ttttgagcaa sasaasaasa asaasaa <210> 431 e211> 392 K212> DNA <213> Nomo sapiens <22205 <221> misc_feature <222> (1)...(392) <2230 n = A,T,C or G 64005 431 gaaaattoag aatggataaa aacaaatgaa gtacaaaata titcagatti acatagogat 60 aaacaagaaa gcacttatca ggaggactta caaatggaag tacactctan aaccatcatc 120 tatcalgget aaatgtgaga blagcacage tgtattattt gtacattgca aacacctaga 180 aagagatggg aaacaaaato ocaggagitt tqtgtgtgga gtootgggtt ttooaacaga 240 catcaticca gcattotgag attagggnga tiggggatca ticiggagit ggaaigitca 300 acaaaagtga igtigtiagg taaaatgtac aacticigga tetatgcaga cattgaaggi 360 grantgagte tygettttae tetyetgitt et <210> 432 <2115 397 <212> DNA <213> Homo sapiens <220× <221> misc_feature <222» (1)...(387) <2235 n * A,T,C or G ggtatochta cataatoaaa tatagotyta gtacatgitt toaitggngt agattaccac 60 asatgesagy cascatytyt systeretty tettatrett tegtetatas tactytatty 120 ngtagtoraa gototoggma gtocagorao tyngaaacat gotocottta gattaacoto 180 gtggacneth ttgttgnatt gtetgaactg tagngeeetg tatittgett etgtetgnga 240 attetyttige tietgygyca litteetigny atgesyagga eeaccacaca gatgacagca 300 atctgaattg ntocaatoac agotgogatt aagacatact gaaatcgtac aggacoggga 360 acaacgtata gaacactgga gtoottt <210> 433 <211> 281 c212> DNA <213> Komo sapiens $<220\times$ <221> misc_feature

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<222> {2}...(281)
<223> m = A,T,C or G
<400> 433
ttcaactago anagaanact gottoagggn stgtaaaatg aaaggottos acgoagttat 60
ctgattasag aacactaaga gagggacaag gotagaagoo goaggatgto tacactatag 120
cagginitat tigggitiggi tygaggagii giggaaaada tygagagatii ggogiiggag 180
stegrogtgy stattecton tightaited accegngagy nichologini geseactgy: 240
tmnasaacog mtatacaata atgatagaat aggacacaca t
<210> 434
<211> 484
<2125 DNA
<213× Nomo sapiens
<400× 434
tittamaata aqoatitaqt qotoaqtooc tactqaqtac tottictoto coctoolotq 60
aatttaatto tticaactig caattigoaa ggattacaca tticacigig atgiatatig 120
tyttycassa asassaasyt ytotttyttt assattactt yytttytysa tocatottyc 180
tttttcccca tiggaactag tdattaaccc atctctgaac tggtagaaa acatctgaag 240
agriagicta icagratoig araggigaat tygaiggito tragaacrat ticarroaga 300
cagootgitt ctatootgit taataaatta giilgggito totacatgoa taacsaacco 360
tgctccaatc tgtcacataa aagtctgtga ottgaagttt agtcagcacc occaccaaac 420
thtattitic talgigittt tigcamcata tgagtgittt gamamimaag imcccatgic 480-
fibbba
<210> 435
<211> 424
<212> DMA
<213> Homo sapiens
<400> 435
gagoogates gagaaggtes attictgest teasogtest oottesagga ageocratgt 60
gggtaggttt saatatogsa ggttsttast setetgeste tataagetea aacceassaa 120
cgatogggca agtaaacccc ctccctcgcc gacttoggaa ctggcgagag ttcagogcag 180
atgggcotgt ggggagggg caagatagat gagggggago ggcatggtgc ggggtgaccc 240
ctiggagaga ggaaaaaaggc cacaagaggg gcigocaccg ccactaacgg agaiggcoct 100
ggtagagaco tüügggggüd üggaadotot ggadtoddda tgdiciaact dodacactot 160
gotatoagaa acttaaactt gaggattito totgiittio actogoaata aattoagago 420
2330
<210> 436
<211× 667
<212> DNA
<213> Homo sapiens
x 2200 >
<221> misc_feature
<222× (1)...($67)
<223> n * A,T,C or G
<400> 436
accttegggaa nactotoaca atataaaggg togtagactt tactccaaat tecaaaaagg 60
toctggoodt gtastoctga asgittitocc aaggitagota taasatoott ataagggigo 120
agonicitot ggaalionio igalitoaaa ginicantoi caagitoiig aaaangaggg 190
cagtteetga aaggeaggta tageaactga tetteagaaa gaggaactgt gtgeaceggg 240
atgggotgec agagtaggat aggattocag atgetgacae ettetgggggg saacaaggget 300
goraggittig tratagract rateasagte oggiteaargi otgigsittig aatalaaare 360
```

PCT/US00/309004

```
igitcatgit tataggacto allosagasi titolalato tolitottak atactolosa 420
agttoataat goigotocat goocagoigg gigagitggo caaatooing iggoostgag 480
gattoottta tggggtoagt gggaaaggtg toaatgggac ttoggtotoc atgoogaaac 540
xccaaagtcx caaacttcaa ctccttggct agtacacttc ggtctagcca gaaaaaaagc 800
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tgttgag
<210> 437
<211> 693
<212> DNA
<213 > Homo sapiens
<400> 437
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taaagottoay gttaggaggo igataagott ggaaggaact toagacagot tiitoagaid 180
ataaaagata attottagoo catgiiotto looagagoag accigaaatg acagcacago 240
aggtactect ctattiticac coetetiget telaciotet ggeagteaga ceigigggag 300
gocatgggag aaageagete tetggatgtt tgtacagate atggaetatt etetgtggae 360
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atthgagttt obgtobytot toagtagagg aaachtbigo bottcacach toacabotga 480
acaectaact getgttgete etgaggtggt gaaagacaga tatagagett acagtattta 540
tectatizet aggeactgag ggetgigggg tacertgtgg tgccaaaaca gateetgtit 600
taaggacatg itgottoaga galgiotgia actalolggg ggololgilg gololliaco 660
etgestesty tgetetetty getgassaty acc
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<211> 360
62125 DWA
<213> Homo sapiens
<400× 438
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ttatgcaatg catcatgcta tttcatacct aatgagggag ttocaggaga ttoaaccagg 120
stytttetae acetytygyt tatyseassy seasetyses asyssiette asyssyyayy 190
actgcaagta tatotggtgg agaagaagga cocaaaaaag acctgttotg tcagtgaatg 240
gataatetaa tgigetteta gtaggeacag ggeteecagg ecaggeetea tteteetetg 300
grototaata gtoaataatt gigiagooai goolaloagi aaaaagalii tigagoaaac 360
<210> 439
<211> 431
<212> DNA
<213> Homo sapiens
c220>
<221> misc_feature
<222> (1)...(431)
<223> n = A,T,C or G
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tygocagggo agcaagoott agoottygot tettgittet gotititiic tygotagace 120
gaagtgtact agccaaggag ttgaagtttg tgactttggt gtttcggcat ggagaccgaa 180
giocoatiga caccittocc acigarosca taaaggaato cicatggsca caaggattig 240
gocaactoso cosgotyggo stygsgosgo attatgasot tygsgagagtat atasgasaga 300
gatatagasa attotigaat gagtootata aacatgaaca ggittatati ogaagcacag 350
acyttgaccy gactitgaty agtyctatga caaacotgyc agcccytcga cycygccycy 420
aatttagtag t
```

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<211> 523
<212> DNA
<213> Homo sapiens
<400× 440
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qqatcittig tatitaagga ticigagati tiqcitgagc aggatiagat aaggcigtic 120
tttasatgic igaaalggaa cagatttcaa aaaaaaacoc cacaatctag ggigggaaca 180
aggaaggaaa gatgtgaata ggctgatggg caassascom atttaccomt cagttccago 240
ctteteteaa ggagaggeaa agaaaggaga tacagtggag acatetggaa agttitetee 300
actggaaaac tgctactatc tgttttttata titctgttaa aatatatgag gctacagaac 360
taxaaattax ascotottig tglooctigg tootggaaca titalgitoo titlaasgaa 420
acaaaaatca aactitacag asagaittga tgtatgtaat acatatagca grictigaag 480
tatatatatc atagcaaata agtoatolga tgagaacaag cta
<2105 441
<211> 430
<212> DMA
<213> Homo sapiens
<400> 441
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iqqccaqqqc aqcaaqcctt aqcctiqqct totiqittot qctitttic iqqciaqacc 120
gaagtgtact agocaaggag ttgaagtttg tgactttggt gtttcggcat ggagaccgaa 180
gtoccattqa cacctttocc actgaccoca taaaggaato ctcatggcca caaggatttg 240
900 agamenta antagagament inagangament pagagagament agamenta antagament 300
gatatagaaa attottogaat gagtootata aacatgaaca ggttiatatt ogaagcacag 360
acyttyaccy gactttgaty agtyctatya caaacctyyc agcccytcya cycgyccycy 420
                                                                  430
aatttagtag
<210> 442
<211> 362
<212> DNA
<213> Homo sapiens
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tttootggaa tgacaattat attittaacti tggtggggga aagagttata ggaccacagt 120
ctroaction gatactigta aattaatott ttantgoact tgttttgaco attaagctat 180
atgittagas atggicatit tacggaasaa tiagaasaat toigataata gigcagaata 240
aatgaattaa tyttitacti aattiatati gaastytesa tyacaaatsa aaattettii 300
tgattatitt tigitticat tiaccagaat aasaactaag saliaasagt tigattacag 360
20
<210> 443
<211> 624
<212> DWA
<213> Somo sapiens
<220×
<221> misc_feature
<222> (1)...(624)
<223> n = A,T,C or G
<4005 443
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ttgaaagaat taaatttaga ggaggggaga gaaagagtat ttagtaggga otgagcasta 120
aatgottatt ttaaaaagaaa tgiaaagago agaaagcaat toaggotaco otgotittig 180
tyrtgyctag tarteogytr gytytrayca yrarytyyra ttysaratty castytyysy 240
eccaasecae agasaatggg gigasatigg cesactitet attasetigg ettecigitt 300
tataaaatat tgigaataat atcacctact tcasagggca gitatgaggc ttaaatgaac 360
teacquotec essecutia escategate scateggiqu esgiectetg teictggisc 420
arggtaasca teettattat taasgtease getaasatga atgtgigtge statgetast 480
agtacagaga gagggcactt aaascaacta agggcstgga gggaaggttt cetggaaaga 540
ngatgottgt getgggtoca aatottggto tactatgaco ttggccaaat tatttaaact 600
ttgtccctat ctgctaaaca gato
<210× 444
<2115 425
<212> DMA
<213> Homo sapiens
₹220»
<221> misc feature
<222> (1)...(425)
<2239 n = A,T,C or 0
<400> 444
quadateatt untutigeat telitigagaa taagaagate agtaaatagi teagaagigg 60
gaagottigt ocaggooigi gigigaacoo aalgittiigo tiagaaatag aacaagtaag 120
ttoattgota tagostasos casasttigo stasgiggtg gicagosast cottgasigo 180
tgottaatgt gagaggttgg taasatoott tgtgosacao totaactooc tgaatgtttt 240
gotatactas garctataca terragaraa queraageta artaaaayaa caarcaacea 300
cototgoaat otgocaccto otgotggcag gatttgtttt tgcatootgt gaagagocaa 360
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gtaga
<210> 445
<23.1> 43.4
<212> DWA
<213> Homo sapiens
<220°≥
<221> misc_feature
<2225 (1)...(414)
<223> n * A,T,C or G
<400> 445
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tgaaaltett igeatgigge agaltatigg atglagtite elitaaciag calalaaate 180
taatatitt caaataaata aacaqcaasa tatqqtaqaa ttaccattta qaacattaiq 240
satgasasat tytyteteta qattatytaa essatsaeta Etteetasee attystetti 300
qqatttttat aatootacto acaaatqact aggottotoo totiqtattt tgaagcagig 360
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<210> 446
<211> 631
<212> DNA
<213> Homo sapiens
<220×
<221> misc_feature
<222> (1)...(631)
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<223> n = A, T, C or G
<400> 446
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totgoatgoa tgggaagtgt gagcattota toaatatgoa ggagocatot tgcaggtgtg 120
atgrtggtta tartggasaa cartgigasa aasaggarta ragigtirta targitgtir 190
coggetotige acquitticas talgitotiaa togoasotigi galtiggaaca attoagatty 240
ctgtcatcty tgtggtggtc ctctgcatca caagggccaa actttaggta atagcattgg 300
acigagattt gtasacittic caacciticca ggasatgooc cagaagcasc agaattosca 360
gacagaagca aaataraggg cartaragtt cagaraatac aaraagagcg tooacgaggt 420
taatotaaag ggagcatgtt toacagtggc tggactacog agagctlyga ctacacaata 480
cagtattata gacsasagaa taagacsaga gatctacaca tgttgccttg catttgtggt 540
aatotacace aatgaaaaca tgtactacag etatatttga ttatgtatgg atatatttga 600
aatagtatac attgtottga tgttttttct g
<210> 447
<211× 505
<212> DNA
<213> Homo sapiens
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<221> misc_feature
<222> (1)...($85)
<223> n \times A,T,C or G
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gootottotg gaattootot gatttoaaag totoactoto aagttottga aaacgagggo 180
agttootgaa aggcaggtat agcaactgat ottoagaaag aggaactgig igcaccggga 240
tgggetgeca gagtaggata ggattecaga tgetgaeaee ttetgggggga aacagggetg 300-
ccaggittgi catageacto atcaaagice ggicaacgic igigoticga aistaaacci 360
gttoatgibb ataggactca bicaagaabb tichatatob etitothata taotobecaa 420
gittoalaatg olgotooatg occagolyyy lyayttyyoo aaaloottyl yyoostyayy 480
atteenttat ggggteagtg ggaaaggtgt caatgggact teggteteea tgcegaaaca 540
ccaaagtcac aaacttcaac toottggcta gtacacttog gtota
<220> 448
<231> 93
<2112> DNA
<213> Somo sapiens
<220×
«ZZl» misc feature
<222> (1)...(93)
<223> n = A,T,C or G
K4005 448
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ggeteeetag tgeeetggag aggangggge tag
<210> 449
<211> 706
<212> DNA
<213> Homo sapiens
<220×
<221> misc_feature
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<222> (1)...(706) <223> n = A,T,C or G <400> 449 ccaagttcat gothtgtgot ggacgotgga caggggggaa aagcnnttgo togtgggtoa 60 ttotgangao ogsacigado atgodagodo tgodgatggt totocatggo toddagtgo 130 cotagaaaq aggtetotag toagagagta gtootagaaq giggcotote ngaggagoca 180 oggggacage atoptgeaga tegtogggeg ogtopcatto gecattoagg otgegeaact 240 gtigggasgg grgatoggig ogggoetett ogetattaog coagetiggog aaagggggat 300 gtgetgesag gegattaagt tgggtaaege cagggtttte ceagtenega egttgtaaaa 360 cgacggccag tgaattgaat ttaggtgacn ctatagaaga gctatgacgt cgcatgcacg 420 cytacgtaag ottggatoot clagagogge ogcotactat tactaaatto yoggoogogt 480 cgacgtggga tecneactga gagagtggag agtgacatgt getggacnet gtecatgaag 540 caetgageag aagetggagg caeaacgene cagacaetca cagetaetca ggaggetgag 600 sacaggitiga accingggagg iggaggitige aaigagetiga gateaggeen eigeneeeca 660 gcatggatga cagagtgasa ctccatctta asassassas asassa <210> 450 <211> 493 <212> DWA <213> Homo sapiens <400> 450 gagacggagt gtcactotgt tgcccaggct ggagtgcagc aagacactgt ctaagaaaaa 60 acagittiaa aaggtaaaac aacataaaaa gaaatatoot atagiggaaa taagagagic 129 asatgagget gagaacitta casagggate thacagacat giogociata teacigcaig 180agrotaagta taagaacaac ottiggggag asarcatost tigacagiga ggiacaatic 240congreaget agigaaatgg giggaattaa acteasatta atectgeeag eigaaaegea 300agagacactg toagagagtt aaaaagtgag tictaloost gaggtgatto cacagtoito 360 teaagteaac acatetgiga acteacagae caagttetta aaccactgit caaacteige 420 tacacatoag aatoacotgg agagotttac aaactoocat tgoogagggt cgacgoggec 480gcgaatttag tag €210> 451 x211% S01 <212> DWA <213 > Homo sapiens <220× <221> misc feature <222> (1)...(501) $\langle 223 \rangle$ n = A.T.C or G <400× 45% gggegegter cattegerat traggetgeg caartgttgg gaagggegat oggtgegge 60 ctottogota thacqccago tqqcqaaaqq ggqatqtqct qcaagqcqat taaqttqqqt 120 aacgccaggg ttttcccagt cocyacgitg taasacgacg gccagtgaat tgaatttagg 180 tqacnctata gaaqaqctat qacqtcqcat qcacqcqtac qtaaqcttqq atcctctaga 240 goggoogoot actactacta aattogoggo ogogtogaog tgggatoone actgagagag 300 tggagagtga catgtgctgg acxctgtoca tgaagcactg agcagaagct ggaggcacaa 380 ogondoagac actcacagot actcaggagg etgagaacag gttgaaacotg ggaggtggag 620 gttgcaatga getgagatea ggeenetgen occcageatg gatgacagag tgaaactoca 480 tottaaaaas saassassas a 501 <230> 452 <2313 51 <212> DNA <213> Homo sapiens

PC1/US00/30904

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<221> misc_feature
<222> {1}...(51)
<223> n = A,T,C or G
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agacggttic accentiacas encettitag gatgggmentt ggggagcaag e
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e210> 453
<211> 317
<212> DNA
<213> Nomo sapiens
<220×
*221> misc_festure
<222> (1)...(327)
<223> n = A,T,C or G
<400× 453
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acatotgaag agotagtota toagoatotg gosagtgast tggatggtto toagaasoat 120
theacceana cagestighth charactight taatasatta ghithiggilic tetacatigea 180
taacaaaroo igotooaalo igicacataa aagicigiga oligaagiit anicagcaco 240
cocaccasar ittatittic taigtgitti tigcaacata igagigtiti gasaataagg 300
tacccatgto tttatta
<210> 454
<211> 231
<212> DNA
<213> Homo sapiens
<400> 454
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teagoracgo cargototty aaggagtott gastbotoot otgotoacto agtagaacca 120
agaagaccaa attottotgo atoocagott gcaaacsaaa tigitottot aggiotocac 180
cottectitt teagigites saagsteets acaatiteat gaasaacage t
<210> 455
<211> 231
<212> DNA
<213> Homo sapiens
<400× 455
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cattyttocy satgygottt coacagycta cacacacacasa acagyasaca tyccaagitt 130
gtttcaacge attgatgact tetecaagga tetteetttg geategacea catteagggg 180
casagasttt otostagoac agetoscast acagggetce titeteetet a
<210× 456
<211> 231
<212> DNA
<213> Homo sapiens
<400> 456
ttggcaggta conttacasa gasgacacca taccitatgo gitattaggi ggastaatca 60
ttocattcag tattatogtt attattottg gagaaaccct gtotgtttac tgtaaccttt 120
tgcactcaaa ttootttato aggaataact acatagocac tatttacaaa gccattggaa 180
```

```
cottettatt tggtgcaget gctagtcagt coctgactga cattgccaag t
                                                           231
<210> 457
<2115 231
<212> DMA
<213> Homo sapiens
<220×
<221> misc feature
<222> (1)...(231)
<223> n = A,T,C or G
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gcattoctta atatgatoti gctataaita gatititoto cattagagtt catacagttt 120
tattigatti lattagoaat oloittoaga agacootiga galcattaag ottigtatoo 180
agrigiciaa alogaigeet californit gaggigiege iggettitgi g
                                                               233
<210× 458
<211> 231
<212> DNA
<213> Homo sapiens
<400> 458
aggicingti occoccasti coactoccci ciactototo tangaetysy cinggoceas 60
agaagagggg tggttagggga agoogttgag acotgaagco coaccotota cottoottca 120
acarrottaac riiggytaac agrattigya attatoatti gygatyayta gaattircaa 180
ggtootgggt taggoatttt ggggggcoag accocaggag aagaagatto t
<210> 459
<211> 231
<212> DWA
<213> Nomo sapiens
<400> 459
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cottogogaa acciqitgqiq qoccascaqt cotaacqqqa caqqacaqaq agacaqaqca 120
goodigaact gitticoote cassacages atootgices teatiggets igigetities 188
actatacaca gtoacogtoc caatgagaaa caagaaggag caccotocac a
<210> 460
<211> 231
<212> DNA
<213> Homo sapiens
<400> 460
graggtataa catgotgcaa caacagalgt gactaggaac ggccggtgac atggggaggg 60
octatoacco taltottiggy ggotgottot toacagtgat catgaagcot agcagcaaat 120
cccacctccc cacacgcaca oggccagcct ggagcccaca gaagggtoct cotgcagcca 180
giggagetig giccageete cagiceacee claceagget taaggataga a
<210> 461
<211> 231
<212> DWA
<213> Homo sapiens
<400> 461
ogaggittiga gaagetotaa tytigeaggyg ageegaag eaggeggeet agggaggyte 60
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gtygggttoa gtyaggagty gyasattygt toagoagaac caagoogtty gytyaataay 188
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<210> 462
<211> 231
<212> 088A
<213> Nomo sapiens
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gaagaactgt tagagagacc aacagggtag tgggttagag atttccagag tottacattt 180
totagaggag gtatttaatt tottotoact catocagtgt tgtatttagg a
<230> 463
<211> 231
<212> DNA
<213> Homo sapiens
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actgagtaga caggigicot otiggoalgg taagiotiaa giococlocc agaiotgiga 120
catttgacag gtgtcttttc ctctggacct cggtgtcccc atctgagtga gasaaggcag 180
tggggaggtg gatottocag togaageggt atagaagece gtgtgasaag e
x210> 464
<2112 231
<212> DNA
<213> Homo sapiene
<400> 464
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congetterag tgacigitging congresses cagetactery graytering tgangueragy 180
ggtgccagog caccagetag atgetetgta acttetagge cecattites e
<210> 455
<211> 231
<212> DNA
<213> Homo sapiens
<400> 465
catgitging tagingtagi aatgoigget gesteteags caggginase thrageheet 60
gtggcaaatt agcaacasat totgacatos tatitatggt tiotgtatot tightgatga 120
aggatggeac aaitittget tgtgiteata alataeteag attagiteag elecaleaga 185
taaactggag acatgcagga cattagggta gtgttgtagc totggtaatg a
<230> 466
<211> 231
<212> DNA
<213> Homo sapiens
<400> 466
caggiacote ittecatigg stactgiget ageasgeatg eteteogygg tittitiast 60
gyvottogaa cagaastigo cacataroca yytataatay titotaacat tigoocayya 120
octytycaat casatattyt yyagaattoo etayotyyay aayteacasa yactatayyo 180
aataatggag accagtocca caagatgaca accagtogtt gtgtgcggct g
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<210> 467
<211> 311
<212> DWA
<213> Homo sapiens
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tgtgccttaa cagaaggtot tgagattota agtgggaato alticagtga cigtcalgtg 180
gcatgggtot otgoocaago togtaatgag actatagcaa ggoggotgtg ggacgttaagt 240
tytyaostys tysysotoos aatagastaa sayysaytys sayttyyass saagayaaya 300
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<230> 468
<231> 3112
<232> DWA
<213> Homo sapiens
<400> 458
cattgtgttg ggagaaaaac agaggggaga tttgtgtggc tgcagccgag ggagaccagg 60
dagatrigca iggigggaag gaccigatga tacagagitt gataggagac aattaaaggc 120
tyyssygdac tygatyccty stystyssyt gysctttcas actygygdac tectyssacy 180
aigggaiggo cagagacaca ggagaigagi iggagcaago toaalaacaa agiggitcaa 240
ogaggactty gasttycsty gagotygago tyaagittay occaattytt tactagitya 300
gtgaatgtgg atgattggat gatcattlet calectetgag cettaggtte cetatecata 360
aaatgyyata caragtatga totataaagt gygatatagt algatotact losotgygtt 430
attigaagga tgaatigaga taatttatit caggigoota gaacaatgoo cagattagta 480
cattigging aactgagaaa tggcataaca ccaaaittaa tatatgicag aigttactat 540
gattateatt caateteata gttttgteat ggeceaattt atecteaett gtgeeteaac 600
aaattgaact gttaacaaag gaatctctgg tootgggtaa tggctgagca ccaotgagea 860
titiosatice agtiggetic tigggtitge tageigeate actagicate tiaaataaat 720-
gaagtiilas caliticios gigatiliit talcicacci tigaagatac talgitaigi 780-
gattasataa agsactigag sagaacaggi ticaliasac alaasatcaa igiagacgca 840
aattttotgg atgggcasta oftstyttos caggasatgo tttaaaatat gcagasgata 900-
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His His His Thr His Glu His Thr Asp Thr Leu Pro Tyr Gly His Trp

His Thr His Cys Bis Thr Val Thr Trp Thr His Leu His Thr Ile Thr

Pro Pro His Thr Leu Pro Val Asp Thr Arg Thr His Arg His Cys His

Thr Asp Thr Gln Asp Thr Val Thr Arg Arg His His His Ala Asp Thr

Pro Pro Leu Trp Cys Arg Leu Asn Tyr Pro Ala Gly Gly Thr Ala Val 130

Als Tyr Ser Cys Leu Ser Asp Trp Leu Ser Pro Gin 135

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Ser His Gly His Thr Gly Ile Val Thr Trp Thr Asp Thr Gln Thr Tyr

Gly Glu Ile Thr Trp Thr His His His Thr Ile Thr Gly Thr Gln Thr

His Gly Asp Ile Thr Thr Trp Thr His Cys His Thr Thr Thr Gly Thr

Arg Asp Ile Thr Leu Ser Sis Cly His Thr Ile Thr His Met Asn Thr

Pro Thr His Cys His Met Asp Thr Gly Thr His Thr Ala Thr Leu Ser 30

Mis Oly Mis Thr Ser Thr Pro Ser Mis Mis Mis Thr Mis Cys Lew Trp 100 1.05

Thr Gln Gly His Thr Asp Thr Val Thr Gln Ile His Lys Thr Leu Ser 120

Mis Gly Asp Ile Thr Met Gln Ile His His His Ser Gly Ala Val 135

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Gly Glu Ile Thr Leu Thr His His His Thr Ile Thr Gly Thr Gln Thr

His Gly Asp Ile The The Tep The His Cys His The The The Gly The

Arg Asp Ile Thr Leu Ser His Gly His Thr Ile Thr His Met Asn Thr

Pro Thr His Cys His Met Asp Thr Ala Thr His Thr Ala Thr Leu Ser

His Gly His Thr Ser Ile Pro Ser His His His Thr His Cys His Val 108

Asp Thr Arg Thr His Arg His Cys His Thr Asp Thr Gln Asn Thr Val

The Arg Arg His His His Ala Asp The Pro Pro His Gly His See The

Arg His Ser Ala Thr Gln Ile His His His Thr Glu Met Arg Thr His 2.50 2.55

Cys His Thr Asp Thr Thr Thr Ser Leu Pro His Phe His Val Ser Ala

Gly Gly Val Gly Pro Thr Thr Leu Gly Ser Asn Arg Glu Ile Thr Trp 283

Thr Tyr Ser Giu Gly Lys Ile She Phe Tyr Phe Leu Gly Asn Cln Ala 200

Arg Leu Cys Leu Lys Lys Arg Lys Lys Lys Gin Tyr Thr Val 238

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Cys Cys Leu Trp Gly Leu Gln Ser Leu Pro Gln Gly Ser Tyr Val Thr 20 30

Val Oly Phe Leu Val Val Lyz Arg Cin Thr lie Gly Arg Leu Glu Arg 35 40

Asp Phe Met The Lyn Cys Ary Lys Gln Pro Gly Leu Pro Pro Ser Gly 50 60

Leu Cys Leu Leu Trp Pro Trp Pro Asn Leu Glu Fhe Gly Arg Arg Gln 65 70 75 80

Asp Arg Leu Thr Trp Ser Ser Val Ser Val Ala Gly Val Cys Ala Cys 85 90 95

Arg Ala Arg Pro Gly Trp Leu Gly Glu Gln Pro Ala Thr Ser Ala Gly
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Ala Gly Vsl Ala Arg Phe Pro Arg Pro Glu Trp Vsl Pro Pro Aso Gly 130 140

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Fro Pro Thr Leu Pro Ser Gln Gly Ser Gly Trp Pro Cys Ser His Ser

Leu Ser Gly Cys His Leu Met Ala Asp Gly Ala Lys Ala Leu Gly Lys 50 55

Als Asp Gly Pro Trp Pro Tyr Leu Phe Val Arg Arg Thr Asp Val Pro

168

63 70 75 Cys Pro Ala Ala Ser Glu Val Gly Gly Cys Ala Pro Ser Ser Trp Arg Als beu Ala Glu Val Thr Gly Cys Ser Leu Gly Pro Leu Gly Leu Ala Gln His Ala Gin Ala Ser Val Leu Leu Leu Cys Tyr Lys Trp Ser His Ile Gly Glu Thr Ser Ser His Leu Arg Ser Lys Val Tyr Ala Ala Phe Gly Gly Ser Ser Pro Cys Leu Lys Gly Leu Met Ser Leu Trp Ala Ser Trp Leu Ser Arg Gly Arg Pro <210> 482 <211> 143 <212> PRT <213> Homo sapiens <400> 482 Net Glu Pro Tyr Arg Gly Asn Lys Lys Gln Val Gin Glu Lys Gly Val Pro Cys Leu Trp Gly Ser Ser Pro Cys Leu Arg Cys His Met Ala Leu Arg Ala Ser Trp Leu Pro Gly Gly Gly Pro Gln Ala Ile Leu Gly Arg Thr Leu Cys Ser Ser Ala Glu Ser Ser Gin Asp Cys His Pro Gly Gly Pro Ser Ile Ala Leu Ala Lys Pro Cys Arg Gly Val Trp Leu Leu Phe Clu Pro Ala Trp Pro Pro Trp His Ala Arg Ala Pro Gly Ala Gly Thr Leu Leu Arg Val Cys Leu Ser Cys Leu Gly Cys His Leu Cys Gly Gly 100 105 Ala Ser Gly Gly Gly Pro Ala Thr Asn Leu Thr Gln Ser Arg Lys

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135

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169

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Gly Phe Leu Val Ala Lys Arg Arg Thr Thr Gly Leu Leu Glu Glu Asp 35 40

Phe Thr Phe Lys Cys Arg Lys Gin Pro Lys Leu Pro Ser Met Arg Leu So S5 60

Ser Leu Leu Trp Pro Trp Arg Asp Leu Lys Phe Val Pro Arg Gln Asp 65 76 80

Lys Leu Thr Arg Ser Ser Val Ser Val Ala Gly Ala Tyr Ala Cys Arg 85 90 95

Ala Cly Pro Cly Trp Leu Lys Clu Cln Pro Ala Thr Ser Ala Arg Vel 100 195 110

Arg Leu Vel Gln Ala Glu His Pro Pro Pro His Pro Leu Glu Glu Val 115 120 125

Gly Met Ala Arg Phe Pro Gin Pro Glu Cys Leu Pro Pro Tyr Cys 130 140

<210> 484

<211× 30

<2125 PRT

<213> Homo Sapien

<400> 484

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33

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170

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Ser Val Ala
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Leu Ala Ser Leu
          20
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     \times 223 \times Made in a lab
     ×400× 493
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Lys Tyr Arg Oly
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     <211> 20
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     <223> Made in a lab
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Leu Met Ile Ser
     <210> 495
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1.72

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3.
Phe Pro Asn Gly
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3 5
Pro Pro Pro Pro Ala
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     <211> 20
     *212> PRT
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i
Ser Val Arg Val
          20
     <210> 498
     <211> 20
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     <223> Made in a lab
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Val Pro Gly Arg
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     <211> 20
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                   5
                                       2.0
Gly Ser Ile Val
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3.
                                                               1.5
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      <211> 414
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                                                                               350
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      <212> DMA
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174

PC1/US00/30904

467

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      <211> 407
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      <400> 506
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175

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  gessaaggee ggttescest eteraassee tegaceareg tggatetess gategeragt
                                                                                                                                                                         300
  organisating aggarange canneather teterages enterested of the contraction of the contract
                                                                                                                                                                         360
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                                                                                                                                                                        420
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178

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Asn Gly Glu Asp Cys Ser Pro His Ser Gln Pro Trp Gln Ala Ala Leu
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                         40
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Val Mot Glu Asn Glu Leu Phe Cys Ser Gly Val Leu Val His Pro Gln 55 ខេច Trp Val Leu Ser Ala Thr His Cys Phe Gln Asn Ser Tyr Thr 11e Gly 70 79 EQ Leu Gly Leu His Ser Leu Glu Ala Asp Gln Glu Pro Gly Ser Gln Met 85 90 Val Glu Ala Ser Leu Ser Val Arg His Pro Glu Tyr Asn Arg Pro Leu 3.00 105 110 Leu Ala Ash Asp Leu Met Leu lie Lys Leu Asp Glu Ser Val Ser Glu 120 3.28 Ser Asp Thr Ile Arg Ser Ile Ser Ile Ala Ser Gln Cys Pro Thr Ala 135 146 Gly Asn Ser Cys Leu Val Ser Cly Trp Gly Leu Leu Ala Asn Gly Arg 3.50 155 Met Pro Thr Val Leu Gln Cys Val Asn Val Ser Val Val Ser Glu Glu 170 165 Val Cys Ser Lys Leu Tyr Asp Pro Leu Tyr His Pro Ser Met Phe Cys 1.80 185 Ala Cly Gly Gly Gln Xas Gln Xas Asp Ser Cys Asa Gly Asp Ser Gly 300 Gly Pro Leu Ile Cys Asn Gly Tyr Leu Gin Gly Leu Val Ser Phe Gly 225 220 Lys Ala Pro Cys Gly Gin Val Gly Val Pro Gly Val Tyr Thr Asn Leu 230 235 Cys Lys Phe Thr Glu Trp Ile Glu Lys Thr Val Gin Ala Ser 248 250 <210> 524 <2115 765 c212> DNA <213> Romo sapien <400> 524 atggccacag caggasatco otggggctgg ttootggggt acctcatoot tggtgtcgca <60 € ggatogotog tototoggtag otgoagocaa atoataaaog googaggactg cagocogoac 120 togoagocot ggcaggoggo actggtoatg gaaaacgaat tgttotgoto gggogtootg 380 gtgcatecge agtgggtget gtcagecgca caetgliter agaacteeta caecaloggg 249 ctgggccige acagiettga ggsegaceaa gageeaggga gecagaiggi ggaggeeage 300 ototoogtac ggcacccaga gtacaacaga cocttgotog ctaacgacct catgetcate 360 aagtiggaog aatoogigto ogagtotgan accatoogga goatcagoat tystiogoag 420 tgorotaccy oggggaacto ttgoctoytt totggotggg gtetgotggo gaacygoaga 483 atgectaccy Egytgeagty cytgaacyty toygtygtyt otgaggagyt otgeagtaag \$40 statatgass ogstytassa osssagsatg ttotgogssg geggagggsa agassagaag 600 gactionigea acggigacie iggggggcon digatoligea acggglacit gcagggeelt 660 gigtoittog gaaaagoose gigiggeeaa giiggegige caggigiota caccaasoic 720 tgcaaartca ctgagtggat agagaaaacc gtccaggcca gttaa 785 <210> 525 <211> 254 <212> PRT <213> Homo sapien <4000 S25 Met Ala Thr Ala Gly Asn Pro Trp Gly Trp Phe Leu Gly Tyr Leu Ile 2.8

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28

3.0

20

180

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3.30

168

184

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25

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The Val Gly Arg Thr Gly Ala Gly Lys Ser Ser Leu Ile Ser Ala Leu 1079 1985

Fhe Arg Leu Ser Glu Pro Glu Gly Lys Ile Trp Ile Asp Lys Ile Leu 1090 1095 1100

Thr Thr Glu Ile Gly Leu His Asp Leu Arg Lys Lys Met Ser Ile Ile 1105 1110 1125

Fro Gln Glu Pro Val Leu Phe Thr Gly Thr Met Arg Lys Asn Leu Asp 1135 1135

Pro Phe Asn Glu His Thr Asp Glu Glu Leu Trp Asn Ala Leu Gln Glu 1140 1150

Val Gln Leu Lys Glu Thr Ile Glu Asp Leu Pro Gly Lys Met Asp Thr 1155 1160 1165

Glu Leu Ala Glu Ser Gly Ser Asn Phe Ser Val Gly Gln Arg Gln Leu 1170 1175 1180

Val Cys Leu Als Arg Als The Leu Arg Lys Asn Ghn The Leu The The 1185 1190 1190 1200

Asp Glu Als Thr Als Asn Val Asp Pro Arg Thr Asp Glu Leu Ile Glu 1219

Lys Lys Ser Gly Arg Asn Leu Pro Thr Ala Pro Cys 1220 1225

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Gln Lys Pro Ser Leu Thr Arg Ala Ile Ile Lys Cys Tyr Trp Lys Ser

lår	Leu 80	Val	Let	Gly	Ile	Phe 55	Thr	Leu	ïle	Gla	Glu 60	Ser	Ala	Lys	Val
110 65	Gln	Pro	Ile	Phe	Leu 70	Gly	Lys	Ile	Ile	Asn 75	Tyr	Phe	Glu	Asn	Tyr 80
Asp	Pro	Met	Asp	Ser 85	Val	Ala	Leu	Asn	Thr 90	Ala	ïyr	Ala	Tyr	Ala 95	The
Val	Leu	Thr	Phe 100	Сув	Thr	Leu	Ile	Leu 105	Ala	Ile	Leu	nis	His 110	Leu	Tyr
Phe	Tyr	His 115	Val	Gln	Cys	Ala	01y 120	Met	Arg	Leu	Ytá	Val 125	Ala	Met	CAs
His	Met 130	Ilæ	Tyr	Ārg	Lys	Ala 135	Leu	Arg	Leu	Ser	Asn 140	Met	Ala	Met	Gly
Lys 145	Thr	Thr	Thr	Gly	Gln 150	Ile	Val	Asn	Leu	Leu 155	Ser	Asn	Asp	Val	Asn 160
Lys	Phe	Assp	Gln	Val 165	Thr	Val	Phe	Leu Leu	His 170	Phe	rea	Trp	Ala	Gly 175	Pro
Leu	Gla	Ala	180 11e	Ala	Val	Thx	Alæ	100 105	Leu	Trp	Met	Glu	Il® 190	Gly	lle
Ser	Cys	Leu 195	Ala	©ly	Met	Ala	Val 200	Leu	Ile	ile	Leu	Leu 205	Pro	Leu	Gln
Ser	Cys 210	Fhe	Gly	Lys	Leu	Phe 215	Ser	Ser	Leu	yığ	Ser 220	Lys	Thr	Ala	Thr
Phe 225	Thr	Asp	Ala	Arg	11e 230	Arg	Thr	Met	Asn	Glu 235	Val	Ile	Thr	Gly	Ile 240
Arg	Ile	Ile	Lys	Met 245	Tyr	Als	Trp	Slu	Lys 250	Ser	Phe	Ser	Asn	Leu 255	Ile
Thr	Asn	Leu	Arg 260	Lys	Lys	Glu	Ile	80x 265	Lys	Ile	Leu	Arg	8er 270	Ser	Cys
Leu	Arg	Gly 275	Met	Asn	Leu	Ala	Ser 280	Phe	Phe	Ser	Ala	Ser 285	Lys	ile	Ile
Val	Ph e 290	Val	Thr	Phe	Thr	Thr 295	Tyr	Val	Leu	Leu	300 300	Ser	Val	Ile	Thr
Ala 305	Ser	Arg	Val	Phe	Val 310	Ala	Val	Thr	Leu	Tyr 315	Gly	Ala	Val	Arg	320 Leu
Thr	Val	The	Leu	Phe 325	Phe	Pro	Ser	Ala	71e 330	3lu	Arg	Val	Ser	91u 335	Ala
Tle	Val	Ser	11e 340	Ārg	årg	Ile	Gln	Thr 345	Phe	Leu	Leu	Leu	Asp 350	Glu	Ile

Ser Gln Arg Asn Arg Gln Leu Pro Ser Asp Gly Lys Lys Wet Val His 360 Val Gln Asp Phe Thr Ala Phe Trp Asp Lys Ala Ser Glu Thr Pro Thr 338 Leu Gln Gly Leu Ser Phe Thr Val Arg Pro Gly Glu Leu Leu Ala Val 398 Val Gly Pro Val Gly Ala Gly Lys Ser Ser Leu Leu Ser Als Val Leu **\$10** Gly Glu Leu Ala Pro Ser His Gly Leu Val Ser Val His Gly Arg Ile 42% Ala Tyr Val Ser Gln Gln Pro Trp Val Phe Ser Gly Thr Leu Arg Ser 440 Asn Ile Leu Phe Gly Lys Lys Tyr Glu Lys Glu Arg Tyr Glu Lys Val He Lys Ala Cys Ala Leu Lys Lys Asp Leu Gin Leu Leu Giu Asp Gly Asp Leu Thr Val Ile Gly Asp Arg Gly Thr Thr Leu Ser Gly Gly Gln Lys Ala Arg Val Asn Leu Ala Arg Ala Val Tyr Gln Asp Ala Asp Ile Tyr Leu Leu Asp Asp Fro Leu Ser Ala Val Asp Ala Glu Val Ser Arg 520 Nis Leu Phe Giu Leu Cys Ile Cys Gln Ile Leu Bis Glu Lys Ile Thr Nr. Ile Leu Val Thr His Gln Leu Gin Tyr Leu Lys Ala Ala Ser Gin Ile 555 Leu Ile Leu Lys Asp Gly Lys Met Val Gln Lys Gly Thr Tyr Thr Glu The Leu Lys Ser Gly Ile Asp The Gly Ser Leu Leu Lys Lys Asp Asn 585 Glu Glu Ser Glu Gln Pro Pro Val Pro Gly Thr Pro Thr Leu Arg Asn Arg Thr Phe Ser Glu Ser Ser Val Trp Ser Sln Glz Ser Ser Arg Pro \$15 Ser Leu Lys Asp Gly Ala Leu Glu Ser Gln Asp Thr Glu Asn Val Pro Val Thr Leu Ser Glu Glo Asn Arg Ser Glu Gly Lys Val Gly Phe Gin

Ala Tyr Lys Asn Tyr Fhe Arg Ala Gly Ala His Trp Ile Val Phe Ile

668 660 670 Phe Leu Ile Leu Leu Asn Thr Ala Ala Gin Val Ala Tyr Val Leu Gin 880 Asp Trp Trp Leu Ser Tyr Trp Ala Asn Lys Gin Ser Met Leu Asn Val 698 Thr Val Asn Gly Gly Gly Asn Val Thr Glu Lys Leu Asp Leu Asn Trp Tyr Leu Gly Ile Tyr Ser Gly Leu Thr Val Ala Thr Val Leu Phe Gly The Ala Arg Ser Leu Leu Val Phe Tyr Val Leu Val Asn Ser Ser Gin Thr Leu Kis Asn Lys Met The Glu Ser The Leu Lys Ala Pro Val Leu Phe Phe Asp Arg Asn Pro Ile Gly Arg Ile Leu Asn Arg Phe Ser Lys Asp Ile Gly His Leu Asp Asp Leu Leu Pro Leu Thr Fhe Leu Asp Phe Ils Gin Thr Leu Leu Gin Val Val Gly Val Val Ser Val Ala Val Ala 820 Val lie Pro Trp lie Ala lie Pro Leu Val Pro Leu Gly lie lie Phe The Phe Leu Arg Arg Tyr Phe Leu Glu Thr Ser Arg Asp Val Lys Arg 840 Leu Glu Ser Thr Thr Arg Ser Pro Val Phe Ser His Leu Ser Ser Ser Leu Gin Gly Leu Trp Thr Ile Arg Ala Tyr Lys Ala Glu Glu Arg Cys Gln Glu Leu Phe Asp Ala His Gln Asp Leu His Ser Glu Ala Trp Phe Leu Fhe Leu Thr Thr Ser Arg Trp Fhe Ala Val Arg Leu Asp Ala Ile 908 Cys Ala Met Phe Val Ile Ile Val Ala Phe Gly Ser Leu Ile Leu Ala Lys Thr Leu Asp Ala Gly Gln Val Gly Leu Ala Leu Ser Tyr Ala Leu Thr Lea Met Gly Met The Gln Trp Cys Val Arg Gln Ser Ala Glu Val Glu Asn Met Met Ile Ser Val Glu Arg Val Ile Glu Tyr Thr Asp Leu 965 370

Glu Lys Glu Ala Pro Trp Glu Tyr Gln Lys Arg Pro Pro Pro Ala Trp 988 985

Pro His Glu Gly Val Ile Ile Phe Asp Asn Val Asn Phe Met Tyr Ser 1000

Pro Gly Gly Pro Leo Val Leo Lys Mis Leo Thr Ala Leo Ile Lys Ser 1010 1015

Gin Glu Lys Val Gly Ile Val Gly Arg Thr Gly Ala Gly Lys Ser Ser 2030 1035

Let Ile Ser Ala Let Phe Arg Let Ser Glu Pro Glu Gly Lys Ile Trp 2050

Ile Asp Lys Ile Leu Thr Thr Glu Ile Gly Leu His Asp Leu Arg Lys 1065

Lys Met Ser Ile Ile Pro Gln Glu Pro Val Leu Phe Thr Gly Thr Met 1080

Arg Lys Asn Leu Asp Pro Phe Asn Glu His Thr Asp Glu Glu Leu Trp 1095

Asn Als Leu Gin Glu Val Gin Leu Lys Glu Thr Tie Glu Asp Leu Pro 1105 1110 1115

Gly Lye Met Asp Thr Glu Leu Ala Glu Ser Gly Ser Asn Phe Ser Val 2125 1130

Gly Gln Arg Gln Leu Val Cys Leu Als Arg Ala Ile Leu Arg Lys Asn 1340 1145

Gln lie Leu Ile lie Asp Glu Ala Thr Ala Asn Val Asp Pro Arg Thr 1160

Asp Glu Leu Ile Gln Lys Lys Ile Arg Glu Lys Phe Ala His Cys Thr 1175

Val Lou Thr Ile Ala His Arg Lou Ash Thr Ile Ile Asp Ser Asp Lys 1190 1195

Ile Met Val Leu Asp Ser Gly Arg Leu Lys Glu Tyr Asp Glu Fro Tyr 1205 1210

Val Leu Leu Gin Asn Lys Glu Ser Leu Phe Tyr Lys Met Val Gin Gin 1225

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199

5 10 35

Met Thr

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<212> PRT

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Ser Val

×210> 546

<231> 29

<212> PRT

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Thr Glu Ala Arg Arg His Tyr Asp Glu Gly Val Arg Met 20

c2105 547

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Ser Ala Pro Ser Leo Ser Pro His Cys Cys Pro Cys Arg Ala Arg Lau 20 25 38

Ala Phe Arg Asn Leu Gly Ala Leu Leu Pro Arg Leu His Gln Leu Cys 35 40 48

Cys Arg Met Pro Arg Thr Leu Arg Arg Leu 50 55

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/AAA\ %49

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Glu Cys

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Gin Ala

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Ser Asp His Trp Arg Gly Arg Tyr Gly Arg Arg Arg Pro Phe

<210> 551

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